

# STIC Search Report Biotech-Chem Library

## STIC Database Tracking Number: 116201

TO: Nita M Minnifield Location: REM-3C18

Art Unit: 1645

Monday, March 08, 2004

Case Serial Number: 09/056019

From: Barb O'Bryen

**Location: Biotech-Chem Library** 

Remsen E01A69

Phone: 571-272-2518

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barbara.obryen@uspto.gov

Search Notes		3.H23.a
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## STIC-Biotech/ChemLib

116201

From:

Chan, Christina

Sent:

Monday, March 08, 2004 8:37 AM

To: Subject: Minnifield, Nita; Pak, Michael; STIC-Biotech/ChemLib

RE: rush sequence search request

#### Please rush. Thanks Chris

Chris Chan

TC 1600 New Hire Training Coordinator and SPE 1644 (571)-272-0841 Remsen, 3E89

----Original Message-----

From:

Minnifield, Nita.

Sent:

Sunday, March 07, 2004 10:43 AM

To:

Chan, Christina

Subject:

rush sequence search request

Christina, please approve, 2 month overdue amdt.

### **STIC**

09/056019

Please do commercial and interference sequence search on SEQ ID NO: 1, 3-5, 7, 9-11, 22-24 (all amino acid sequences).

Please provide a paper copy of results.

Transferred case

Thanks, Nita M. Minnifield

Searcher:
Phone:
Location:
Date Picked Up:
Date Completed:
Searcher Prep/Review:
Clerical:
Online time:

TYPE OF SEARCH:
NA Sequences:
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other:

VENDOR/COST (where applic.) STN:
DIALOG:
Questel/Orbit:
DRLink:
Lexis/Nexis:
Sequence Sys.:
WWW/Internet:
Other (specify):

Art Unit 1645
Office REM-3C01
Mailbox REM-3C18
571-272-0860

Searcher:
Phone:
Location:
Date Picked Up:
Date Completed:
Searcher Prep/Review:
Clerical:
Online time:

TYPE OF SEARCH:	
NA Sequences:	
AA Sequences:	
Structures:	
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Litigation:	
Full text:	
Patent Family:	
Other:	

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DRLink:
Lexis/Nexis:
Sequence Sys.:
WWW/Internet:
Other (specify):

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to the EMBL/GenBank/DDBJ databases.
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Pfan; PF00746; Gram pos anchor; 1.
Pfan; PF04650; RICH; 2.
Pfan; PF04650; VSIRK signal; 1.
PIGRFAMs; TIGRO1169; VSIRK signal; 1.
PROSITE; PS50847; GRAM POS ANCHORING; 1.
SEQUENCE 769 AA; 85266 WW; D96DBAD7C2E5
      Submitted (MAY-1999) to the EMBL/GenBany-i-SUBGELLULAR LOCATION: ATTACHED TO THE AMIDE BOND (BY SIMILARITY).

EMBL; APT-54014; AAR73804.1; -GO; GO:0005618; C:cell wall; IEA.

GO; GO:0016020; C:membrane; IEA.
InterPro; IPR005877; Gpos_YSIRK.
InterPro; IPR001899; Gram pos_anchor.
InterPro; IPR001899; Gram pos_anchor.
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Pfam; PF05062; RICH; 1.
Pfam; PF04650; YSIRK signal; 1.
TIGRFAMs; TIGR01168; YSIRK signal; 1.
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InterPro; IPR002479; CW binding.
InterPro; IPR005877; Gpos YSIRK.
InterPro; IPR007756; RICH.
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NCBI_TaxID=1313;
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Submitted (MAX-1999) to the EMBL/GenBank/DDBJ databases.
-1- SUBCELLULAR LOCATION: ATTACHED TO THE CELL WALL PEPTIDOGLYCAN I AN AMIDE BOND (BY SIMILARITY).
EMBL; ARISTO4043, ARF73816.2;
GO; GO:0005618; C:cell wall; IEA.
GO; GO:0016020; C:membrane; IEA.
InterPro; IPR001899; Gram_Dos_anchor.
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                                                                                                                                                                                                                                                             Tannelli F., Oggioni M.R., Pozzi G., "Allelic Variation in the Highly Polymorphic Locus pspc of
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                                                                                     Streptococcus pneumoniae.
Bacteria, Firmicutes, Lactobacillales, Streptococcaceae,
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Bacteria; Firmicutes; Lactobacillales; Streptococcaceae;
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"Allelic Variation in the Highly Polymorphic Locus pspC
Streptococcus pneumoniae.";
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      Last annotation update)
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Pfam, PF05162; NICH; 2.
Pfam, PF04650; YSIRK signal; 1.
TIGRFAMS; TIGR01167; "IPXTG anchor; 1.
PROSTITE; PS56847; GRAM POS AGTAIL; 1.
PROSTITE; PS56847; GRAM POS AGTAIL; 1.
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01-OCT-2003 (TrEMBLrel. 25,
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Best Loca Matches

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96 KKTVDIVNKLONINNEYLNKIIQSTSTYBELQKLAMBSQSEVDKAVSEFEKDLSSSSSG 155
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STRAIN=serotype 1;
MEDLINE=98010350; PubMed=9350867;
Hammerschmidt S., Talay S.R., Brandtzaeg P., Chhatwal G.S.;
"SpsA, a novel pneumococcal surface protein with specific binding to Immunoglobulin A and secretory component.";
MMO1. Microbiol. 25.1113-1124 (1997).
EMBL; Y10818; CAA71783.1;
TO THE CELL WALL PEPTIDOGLYCAN BY
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Bacteria; Firmicutes; Lactobacillales; Streptococcaceae;
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Last annotation update)
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106 NFALNLKLSRIKTEYLYGLSVLKEKSEAELPSKAELPSKIKEKLTAAFEHFKKOT---- 160
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Surface protein PSpC.
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NCBI_TaxID=1313;
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                                                                                                                                                                   Gaps
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A Tannelli F., Oggioni M.R., Pozzi G.;

"Allelic Variation in the Highly Polymorphic Locus pspC of Streptococcus preumoniae.";

I Submitted (MAY-1599) to the EMBL/GenBank/DDBJ databases.

EMBL, AF154021; AAF73790.1;

R GO; GO:0016020; Camembrane; IEA.

R InterPro; IPR002479; CW binding.

R InterPro; IPR005877; GpGs YSIRK.

R InterPro; IPR005877; GpGs YSIRK.

R InterPro; IPR005837; Tropomyosin.

R Pfam; PF00457; W binding_1; 9.

R Pfam; PF00465; RICH; 1.

R Pfam; PF00465; RICH; 1.

R Pfam; PF004650; YSIRK signal; 1.

R Pfam; PF04650; YSIRK signal; 1.

R PRINTS; PR00194; TROPOMYOSIN.

R TIGRRAMS; TIGR01168; YSIRK signal; 1.

SEQUENCE 709 AA; 80251 WW; 32BBC56E380EBB7A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    56.5%; Score 790; DB 2; Length 709; 58.5%; Pred. No. 4.7e-31; ive 38; Mismatches 38; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Streptococcus pneumoniae.
Barteria, Firmicutes, Lactobacillales; Streptococcaceae, Streptococcus, NCBI_TaxID=1313;
                      1 37 POTENTIAL.
38 523 POTENTIAL.
523 AA, 59151 MW, B3390DF8A706465A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Created)
Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 58.5%
Matches 175; Conservative
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                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     241
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Best Local
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09KK38;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     181 ESDVEVKKAELELVKVKANEPRDEQKIKQABAEVESKQAEATRLKKIKTDREEAEBEAKR 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    240
                                                                                                                     EIAESDVEVKKAELELVKVKANEPRDEQKIKQAEAEVESKQAEATRLKKIKTDREEAEEE 237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   61 TITVALVNELANJIKARFLAKIVESTSESQLQILAMMESRSKVDEAVSKFEKDSSSSSSSS 120
SDSSTKPEASDTAKPNKPTEPGEKVAEAKKKVEEAEKKAKDQKEEDRRNYPTITYKTLEL 177
                                                         39 EXEVITQVPISSNRANKSQ-----TEHMRAAKQVDEXIKGGL-----QLDRRKH 82
                                                                                                                                                                                                                                        238 AKRRADAK------EQGKPKGRAKRGVPGELATPDKKENDAKSSDSSVGEETL 284
                                                                                                                                                                                                                                                                                    1 ENEGATOVPISSNRANESQAEQGEQPKKLDSERDKARKEVEEYVKKIVGESYAKSIKKRH 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              203 BIABSDVBVKKABLELVKBEANKPRNEQKVKQAKAKVBSKQABATRLEKIKTDREBAEBE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             241 RADAK-------EQGKPKGRAKRGVPGELATPDKKENDAKSSDSSVGEBTL 284
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ||:||
| RAEAKLKEAVEKNTATSEQGKPKRRGKRGALGEQATPDKKENDAKSSDSSVGEBTL 296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          50; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=set25;

A lanelli F., Oggioni M.R., Pozzi G.;

I alnelli F., Oggioni M.R., Pozzi G.;

I allello Variation in the Highly Polymorphic Locus pspC of Streptococcus pneumoniae.";

E Submitted (MAY-1999) to the EMBL/GenBank/DDBJ databases.

E MBLG AFT54040; AAF73812.1; -...

R GG: 0016020; C:membrane; IEA.

R InterPro; IPR005479; CW_binding.

R InterPro; IPR007479; CW_binding.

R InterPro; IPR00777; GPOSS YSIRK.

R Pfam; PF01473; CW_binding.1; 7.

R Pfam; PF01650; XIRK signal; 1.

R Pfam; PF05650; XSIRK signal; 1.

R TIGRFAMS; TIGR0168; YSIRK signal; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Streptococcus pneumoniae.
Bacteria; Firmicutes; Lactobacillales; Streptococcaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Last sequence update)
Last annotation update)
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54.6%; Score 763; DB 2;
Best Local Similarity 57.4%; Pred. No. 9.3e-30;
Matches 170; Conservative 35; Mismatches 41;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                             2 EPGEKVAEAKKKVEEAEKKAKDQKEEDRRNYPTITYKTLELEIAESDVEVKKAELELVKV 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 38, Application US/09536784
Patent No. 6573082
GENERAL INFORMATION:
APPLICANT: Choi et. al.
APPLICANT: Choi et. al.
NUMBER OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
SIREET: 9410 Key West Avenue
CITY. Rockville
                                                                                                                                                                                                                                                                                                                                              0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ch 81.5%; Score 433; DB 4; Length 453;
1 Similarity 82.1%; Pred. No. 5.3e-33;
87; Conservative 15; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                            Query Match 81.5%; Score 433; DB 3; Length 453; Best Local Similarity 82.1%; Pred. No. 5.3e-33; Matches 87; Conservative 15; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    148 EAKEPRNEEKVKQAKAEVESKKAEATRLEKIKTDRKKAEEBAKRKA 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         62 KANEPRDEOKIKQAEAEVESKQAEATRLKKIKTDREEAEEBAKRRA 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
COMPUTER: HP VECTRA 486/33
OPERATING SYSTEM: MSDOS Version 6.2
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/536,784
FILING DATE: 30-OCC-1997
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION UNDERS: 06/961,083
FILING DATE: OCT-30-1997
ATTORNEY/AGENT INFORMATION:
NAME: MICHOELLE &: Marks
REGISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB340P3
TELECHANDIN INFORMATION:
TELEPHANE: (301) 309-8504
TELEFAX: (301) 309-8512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear

MOLECULE TYPE: procein

SEQUENCE DESCRIPTION: SEQ ID NO: 38:
US-09-536-784-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 453 amino acids
TYPE: amino acid
STRANDEDNESS: single
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8514
TELEFAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO; 38;
                                                                                                                         LENGTH: 453 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Maryland
COUNTRY: USA
                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: protein
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Best Local Similarity
Matches 87; Conservat
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APPLICANT: Briles, David E.
APPLICANT: McDaniel, Larry S.
APPLICANT: Swiatlo, Edwin
APPLICANT: Yother, Janet
APPLICANT: Crain, Marilyn J.
APPLICANT: Hollingshead, Susan
APPLICANT: Tart, Rebecca
APPLICANT: Tart, Rebecca
APPLICANT: PRODOKS: Walter, Alexis
TITLE OF INVENTION: EXPRESSION PRODUCTS THEREPROM, AND USBS OF SUCH GENES,
TITLE OF INVENTION: PORTIONS AND PRODUCTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88 KPEKKVAEAEKKKVEEAKKKAEDQKEEDRANYPINTYKTLELEIAESDVEVKKAELELVKE 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 EKVARAKKKVERAEKKAKDOKEEDRRNYPTITYKTLELEIAESDVEVKKAELELVKVKAN 64
2 EPGEKVASAKKKVEBAEKKAKDQKEEDRRNYPTITYKTLELBIAESDVEVKKAELELVKV 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4; Gaps
                                                                                                                81.5%; Score 433; DB 4; Length 864; 86.7%; Pred. No. 1.1e-32; tive 8; Mismatches 2; Indels
                                                                                       62 KANEPRDEOKIKOAEAEVESKOAEATRIKKIKTDREEAEEEAKRRA 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               65 EPRDEOKIKOAEAEVESKOAEATRIKKIKTDREEAEEEAKRRADA 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC compatible
COMPUTER: SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATE:
FILING DATE: 16-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SSEE: Curris, Morris & Safford, P.C. 1530 Fifth Avenue New York New York 19. U.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         454312-2460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Frommer Esq., William S. REGISTRATION NUMBER: 25,506
                                                                                                                                                                                                                                                  Sequence 40, Application US/08714741
Patent No. 6500613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 456
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
TELEPAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                864 amino acids
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TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 4.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-714-741-40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
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RESULT 12 US-09-286-981B-10

0; Gaps

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MOLECULE TYPE: amino acid US-08-714-741-46
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
TITLE OF INVENTION:
   TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New York : U.S.
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                           172 KEKVESKKAEATRIJENIKTDRKKA-EBEAKRKADAKLKEANVATSDQGKPKGRAKRGVPG 230
                                                                                                                                                                                                                                                                                                                                                                         61 KYLRELNVLEEKSKOELPSEIKAKLDAAFEKFKKOTL--KPGEKVAEAKKKVE----- 111
                                                                                                                                                                                                                                                                            112 BAKKKAEDOKEEDRRNYPINTYKTLELEIAEFDVKVKEAELELVKEEAKESRNEGTIKQA 171
                                                                                                                                                                                                                                                                                                    115 EAKKKAKAQKEEDHENYPTITYKTLDLEIAEFDVKVKEABLELVKKEADESRNEGTINQA 174
                                                                                                                              1 ENEGSTQAATSSNMAKTEHRKAAKOVVDEYIEKMIREIQLDRRKHTQNVALNIKLSAIKT 60
                                                                                                                                                    2 ENBRITQVPTSSNRGKPERRKAARQ-FDEYINKM---IQLDKRKGTQNLAFNIQLSRIKT 57
                                                                                            Gapa
                                                                                            23;
                                                   Query Match
66.1%; Score 826.5; DB 4; Length 419;
Best Local Similarity 70.1%; Pred. No. 1.1e-57;
Matches 185; Conservative 26; Mismatches 30; Indels 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: PNEUMOCOCCAL GENES, PORTIONS THEREOF
TITLE OF INVENTION: EXPRESSION PRODUCTS THEREFROM, AND U
TITLE OF INVENTION: PORTIONS AND PRODUCTS
NUMBER OF SEQUENCES: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.30
URRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/714,741
FILING DATE: 16-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5: Curtis, Morris & Safford, P.C
530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Frommer Esq., William S.
REGISTRATION NUMBER: 25,506
REPRENCE/DOCKET NUMBER: 454312-2460
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                229 EQATPDKKENDAKSSDSSVGEETL 252
                                                                                                                                                                                                                                                                                                                                                                                                                            231 ELATPDKKENDAKSSDSSVGEETL 254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
; ORGANISM: Streptococcus pneumoniae US-09-286-981B-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Brooks-Walter, Alexis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 46, Application US/08714741
Patent No. 6500613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yother, Janet
Crain, Marilyn J.
Hollingshead, Susan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                McDaniel, Larry S.
Swiatlo, Edwin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Briles, David E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New York
: New York
RY: U.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Briles
APPLICANT: McDani
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 16
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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US-08-714-741-46
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THEREFROM, AND USES OF SUCH GENES,
                                                                                                                                         47 AKTEHRKAAKKVVDEYIEKMLREIQLDRRKHTQNVALNIKLSAIXTKYLRELSVLKENSK 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        227 VKVKEABLELVKBEANESRKXKAXDQKBEDRRNYPTNTXKTLDLEIAEXDVKVKEABLEL 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    287 VKEBAXEXRNEEKIKQAKEKVESKKARATRLEKIKTDRKKABEBAKRKAEESEKKA--AE 344
                                                                                                                                                                                                                                                                                                                                                                      226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             163 -----RNEGTIKQAKEKVESKKAEATRLENIKTDRKKAEBEAKRKADAKLKEANVAT 214
                                                                                                                                                                                                                                                                                                                   85 IDAAFEKFKKDTLKPGEKVAEAKKKVEEAKKKAEDOKEEDRRNYPTNTYKTLELEIAEFD 144
                                                                                                                                                                                                               70 -----EBKSKDELPSEIKAK 84
                                                                                                                                                                                                                                                              107 KEELTSKTKAELTAAFEQFKKDTLKPEKKVAEAEKKVEEAELXVXEEKSXXELPSEIKAK
                                                                                                                                                                                                                                                                                                                                                     167 LDAAFXKFKKDTLKPGEKVAEAKKCVEEAKKKAKDQKEEDRRNYPTNTYKTLELEIAESD
                                                                                                          15 AKTEHRKAAKQVVDEYIERMLREIQLDRRKHTQNVALNIKLSAIKTKYLRELNVL----
                                                     28; Indels 106; Gaps
     Length 605;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/714,741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  215 SDQGKPKGRAKRGVPGELATPDKKEN---DAKSSD 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                345 ADEEKIK-QAKAKVESKKAEATRLENIKTDXKKAE 378
Query Match 64.3%; Score 805; DB 4; Best Local Similarity 56.7%; Pred. No. 8.4e-56; Matches 190; Conservative 11; Mismatches 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3: Curtis, Morris & Safford, P.C.
530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       : PORTIONS AND PRODUCTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Palter, Alexis
PNEUMOCOCCAL GENES,
EXPRESSION PRODUCTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   454312-2460
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MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         :-08-714-741-40
Sequence 40, Application US/08714741
Patent No. 6500613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Briles, David E.
McDaniel, Larry S.
Sviatlo, Edwin
Yother, Janet
Crain, Marilyn J.
Hollingshead, Susan
Tart, Rebecca
Brooks-Walter, Alexis
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                                                                                                                                                                                                                                                                                                                                                                                                                        145 VKVKEAELELVKEEAKES----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             : (212) 840-333
(212) 840-0712
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS: ADDRESS: ADDRESSEE: Curtis, N
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229 TLPSPSLNWANESQTEHRKD----VDEYIKKMLSEIQLDRRKHTQNVNLNIKLSAIKTKY 284
                                                                                                                                                                                                                                                                                                                                                                                                                                  63 IRELNVLEEKS-KDELPSEIKAKLDAAFEKFKKDTLKPGEKVAEAKKKVEBAKKKAEDOK 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  285 LYELSVLKENSKKEELTSKTKAELTAAFEQFKKOTLKPEKKVAEAEKKVEEAKKYEEAKKAKAKOOK 344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   122 BEDRRNYPTNTYKTLELEIAEFDVKVKEAELELVKEEAKESRNEGTIKOAKEKVESKKAE 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      345 EEDRRNYPINTYKTLELEIAESDVKVKEAELELVKEEANESRNEEKIKQAKEKVESKKAE 404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   182 ATRIENIKTDRKKAEEEAKRKADAKUKEANVATSDQGKPKGRAKRGVPGELATPDKK--- 238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ------AEAKOKVDAEEYALEAKIAE 453
                                                                                                                                                                                                                                                                                                                                    6 TOAATSSNWA---KTEHRKAAKQVVDEYIEKMLREIQLDRRKHTQNVALNIKLSAIKTKY
                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                              Query Match 61.6%; Score 770; DB 4; Length 864; Best Local Similarity 67.0%; Pred. No. 7.5e-53; Matches 177; Conservative 18; Mismatches 39; Indels 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      405 ATRLEKIKTDRKKAEEEARRKAEESEKKA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         239 -----ENDAKSSDSSVGEETL 254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 : | | | : | | 454 LEYEVQRLEKELKEIDESDSEDYL 477
information for SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
LENGTH: 864 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: amino acid
US-08-714-741-40
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Search completed: March 8, 2004, 12:37:48 Job time: 11.7102 secs

us-09-056-019b-10.rai

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STRANDEDNESS:
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         ADDRESSEE:
STREET: 530
CITY: New Y
STATE: New
COUNTRY: U.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 KSGKKVAEAEKKVEBAEKKAKDQKEEDRRNYPTNTYKTLDLEIAESDVKVKEAELELVKE 60
                                                 APPLICANT: Choi et. al.
TITLE OF INVENTION: Streptococcus pneumoniae Antigens and Vaccines
NUMBER OF SEQUENCES: 452
CORRESPONDENCE ADDRESS:
ADDRESSEE: Human Genome Sciences, Inc.
STREET: 9410 Key West Avenue
CITY: Rockville
STATE: Maryland
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     148 EAKEPRINEEKVKQAKAEVESKKAEATRLEKIKTDRKKAEEEAKRKA 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 453;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3; Indels
                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb storage COMPUTER: HP Vectra 486/33 OPERAING SYSTEM: MSDOS Version 6.2 SOFTWARE: ASCII Text CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 468; DB 4;
Pred. No. 1.7e-35;
8; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/536,784
FILING DATE: 30-Oct-1997
CLASSIFICATION: <UNKNOWED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/961,083
FILING DATE: 0CT-30-1997
ATTORNEY/AGENT INFORMATION:
NAME: MIChelle S. Marke:
RESISTRATION NUMBER: 41,971
REFERENCE/DOCKET NUMBER: PB340P3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; MOLECULA TYPE: protein
; SEQUENCE DESCRIPTION: SEQ ID NO: 38:
US-09-536-784-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
TELEFAX: (301) 309-8512
Application US/09536784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 453 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 38
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 89.6
Matches 95; Conservative
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CAL GENES, PORTIONS THEREOF,
NA PRODUCTS THEREFROM, AND USES OF SUCH GENES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          348 KPEKKVABAEKKVEBAKKYAKDQKEEDRRNYPINIYKTIELEIAESDVKVKEAELELVKE 407
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 KSGKKVAEAEKKVEEAEKKAKDQKEEDRRNYPINTYKTLDLEIAESDVKVKEAELELVKE
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2.8e-35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 EAKEPRDEEKIKQAKAKVESKKAEATRLENIKTDRKKAEEEAKRKA 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER RELABBLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: END FO Compatible
COMPUTER: END FO Compatible
COMPUTER: END FO Compatible
COMPUTER: END FO COMPATION
COMPATE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/714,741
FILING DATE: 16-SEP-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Frommer ESG., William S.
REFERENCE/DOCKET NUMBER: 454312-2460
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 840-373
INFORMATION FOR SEG ID NO: 42:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 90.3%; Score 467; DE Best Local Similarity 91.5%; Pred. No. 2.8e Matches 97; Conservative 3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Crain, Marilyn J.
APPLICANT: Hollingshead, Susan
APPLICANT: Tart, Rebecca
APPLICANT: Brooks-Waleer, Alexis
TITLE OF INVENTION: EXPRESSION PRODUCTS
TITLE OF INVENTION: EXPRESSION PRODUCTS
NUMBER OF SEQUENCES: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 40, Application US/08714741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
1: Curtis, Morris
530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NOLECULE TYPE: amino acid
US-08-714-741-42
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321 KPEKKVABAEKKVEBAKKKAKDOKBEDRRNYPINTYKTLELEIAESDVKVKEAELELVKE 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 KKVARABEKKVEBAKKKAEDQKEBDRRNYPTNTYKTLELBIAESDVEVKKAELBLVKEBAK 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 KSGKKVABABKKVEBABKKAKDOKBBDRRNYPTNTYKTLDLBIABSDVKVKBABLBLVKB 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 KKVAEAEKKVEEAEKKAKOOKEEDRRNYPTNTYKTLDLEIAESDVKVKEAELELVKEEAK 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Wizemann, Theresa M.
APPLICANT: Wizemann, Theresa M.
APPLICANT: Koenig, Scott
APPLICANT: Koenig, Scott
APPLICANT: Scott
APPLICANT: Scott
APPLICANT: Obneson, belivatives of Choline Binding Proteins for Vaccines
TILE REFERENCE: 469201-396
CURRENT APPLICATION NUMBER: US/09/286,981B
CURRENT FILING DATE: 1999-040-6
PRIOR PLING DATE: 1999-05-15
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 1
LENGTH: 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 89.6%; Score 463; DB 4; Length 103; Best Local Similarity 92.2%; Pred. No. 8.9e-36; Matches 95; Conservative 6; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 90.3%; Score 467; DB 4; Length 864; Best Local Similarity 91.5%; Pred. No. 4.3e-35; Matches 97; Conservative 3; Mismatches 6; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       381 BANESRNEEKIKQAKEKVESKKAEATRIEKIKTDRKKAEEEAKRKA 426
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/714,741
FILING DATE: 16-SEP-1996
CLEASIFICATION: 458-1996
ATTORNEY AGENT INFORMATION:
NAME: Frommer Esq., William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454312-2460
TELEROWAUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEROMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
LENGTH: 6AAACTERISTICS:
LENGTH: 6AAACTERISTICS:
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT (CRGANISM: Streptococcus pneumoniae US-09-286-9818-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/09286981B Patent No. 6503511 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: amino acid
US-08-714-741-40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-286-981B-1
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completed: March me : 5.46962 secs

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(212) 840-0712
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TELEFAX:
                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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APPLICANT: Brides, David E.
APPLICANT: Swallel, Larry S.
APPLICANT: Swallel, Swall
APPLICANT: Crain, Marilyn J.
APPLICANT: Tart, Rebecca
APPLICANT: Brooks-Walter, Alexis
TITLE OF INVENTION: FRERESSION PRODUCTS THEREFROM, AND USES OF SUCH GENES,
TITLE OF INVENTION: PORTIONS AND PRODUCTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100 PGEKVABAKKKVEBAKKKABDQKBEDRRNYPINIYKILELBIAEFDVKVKBAELELVKBE 159
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 PGEKVABAKKKVEEAKKKAEDQKEEDRRNYPTNTYKTLELEIAEFDVKVKBAELELVKEE 60
                                                                                                          APPLICANT: Missing, instruction, APPLICANT: Moreing, APPLICANT: Scott APPLICANT: Johnson, Leslie S
TITLE OF INVENTION: Derivatives of Choline Binding Proteins for Vaccines FILE REFERENCE: 469201-396
CURRENT APPLICATION NUMBER: US/09/286,981B
CURRENT APPLICATION NUMBER: US 60/085,743
PRIOR FILING DATE: 1998-05-15
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 4
LEARLY APPLICATION NUMBER: US 60/085,743
SEQ ID NO 4
LEARLY APPLICATION NUMBER: US 60/085,743
LEARLY APPLICATION NUMBER: US 60/085,743
SOFTWARE: Patentin Ver: 2.1
LENGTH: 251
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPPRATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/714,741
FILING DATE: 16-SEP-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Curtis, Morris & Safford, P.C.
530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: PRT
; ORGANISM: Streptococcus pneumoniae
US-09-286-981B-4
                Sequence 4, Application US/09286981B
Patent No. 6503511
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NAME: Frommer Beg., William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 45431
TELECHONE: (212) 840-3333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 42, Application US/08714741
Patent No. 6500613
                                                                                               Theresa M.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: New York
STATE: New York
COUNTRY: U.S.
ZIP: 10036
                                                                                               Wizemann,
                                                                         GENERAL INFORMATION:
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ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYER: Rloppy disk
COMPUTER: RAPPO disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFFWARE: Percent Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/714,741
FILING DATE: 16-SEP-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Frommer Esq., William S.
REGISTRATION NUMBER: 25,506
REGISTRATION NUMBER: 25,506
RECISTRATION NUMBER: 25,506
RECISTRATION NUMBER: 25,506
TELEPRAK: (212) 840-3133
TELEPRAK: (212) 840-0712
INFORMATION POR SEQ ID NO: 40:
SEQUENCE CHRRACTERISTICS:
TENTAL READAMER: SEQ ID NO: 40:
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APPLICANT: Yother, Janet
APPLICANT: Crain, Marilyn J.
APPLICANT: Crain, Marilyn J.
APPLICANT: Tark, Rebeca
APPLICANT: Tark, Rebeca
APPLICANT: Brooks-Walter, Alexis
TITLE OF INVENTION: EXPRESSION PRODUCTS
TITLE OF INVENTION: EXPRESSION PRODUCTS
TITLE OF INVENTION: PORTIONS AND PRODUCTS
TITLE OF INVENTION: PORTIONS AND PRODUCTS
TITLE OF INVENTION: PORTIONS AND PRODUCTS
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                                                                                                      588 amino acida
                                                                                                                                                                                                                                                   ) TOPOLOGY: linear
) MOLECULE TYPE: amino acid
US-08-714-741-42
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: siz
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RESULT 10
US-09~286-981B-19
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APPLICANT: Kizemann, Theresa M.
APPLICANT: Kornig, Scott E.
APPLICANT: Moderage Scott E.
APPLICANT: Moderage Scott E.
TILLE OF INVENTION: Derivatives of Choline Binding Proteins for Vaccines
TILLE OF INVENTION: Derivatives of Choline Binding Proteins for Vaccines
TILLE OF INVENTION: 4099-04-06
CURRENT APPLICATION NUMBER: US/09/286,9818
CURRENT APPLICATION NUMBER: US/09/286,9818
PRIOR APPLICATION NUMBER: US/09/286,9818
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 15
LENGTH: 419
                                                                                                                              1 PGEKVABAKKKVBBAKKKABDQKBEDRRNYPTNTYKTLBLBIABFDVKVKBABLBLVKBB 60
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APPLICANT: Wizemann, Theresa M.
APPLICANT: Wizemann, Theresa M.
APPLICANT: Woenig, Scott
APPLICANT: Commisson, Lesiis S.
TITLE OF INVENTION: Derivatives of Choline Binding Proteins for Vaccines FILE, REPREMENT: 469201-336
CURRENT FILING DATE: 1990-04-06
PRIOR FILING DATE: 1990-04-06
PRIOR PILING DATE: 1998-05-15
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO. 3
SEX ID SEX
                                           0; Gaps
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86.1%; Score 452; DB 4; Length 103;
Best Local Similarity 91.3%; Pred. No. 1.5e-34;
Matches 94; Conservative 4; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                   -61 AKESANEGTIKOAKEKVESKKAEATRLENIKTORKKAEEEAKRKAD 106
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Best Local Similarity 90.6%; Pred. No. 1.4e-34;
Matches 96; Conservative 4; Mismatches 6; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-286-981B-15; Application US/09286981B; Sequence 15, Application US/09286981B; Patent No. 6503511; Patent No. 650351; APPLICANT; Wizemann, Theresa M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ) TYPE: PRT; ORGANISM: Streptococcus pneumoniae US-09-286-981B-15
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Patent No. 6503511
GENERAL INFORMATION:
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APPLICANT: Wizemann, Theresa M.
APPLICANT: Wizemann, Theresa M.
APPLICANT: Gondson, Leslie S
FILE OF INVENTION: Derivatives of Choline Binding Proteins for Vaccines
FILE REFERENCE: 469201-396
FILE REFERENCE: 469201-396
FILE REFERENCE: 1999-04-06
FILE REFERENCE: 1999-04-06
FILE REFERENCE: 1999-04-06
FILE REFERENCE: 1999-05-18
NUMBER OF SEQ ID NOS: 38
SOFTWARE: Patentin Ver. 2.1
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                                   APPLICANT: Wisemann, Theresa M.
APPLICANT: Wisemann, Theresa M.
APPLICANT: Soenig, Scott
APPLICANT: Colonson, Lesile S
TITLE OF INVENTION: Derivatives of Choline Binding Proteins for Vaccines
FILE REFERENCE: 469201-396
FILE REFERENCE: 469201-396
CURRENT APPLICATION NUMBER: US 60/085,743
PRIOR APPLICATION NUMBER: US 60/085,743
PRIOR APPLICATION NUMBER: US 60/085,743
SOFTWARE: PALENTING DATE: 1999-05-15
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PALENTIN Ver. 2.1
3 EKVABAKKKVEBAKKKAEDQKEEDRRNYPINTYKTLELEIAEFDVKVKEAELELVKEBAK 62
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
86.1%; Score 452; DB 4; Length 114;
Best Local Similarity 91.3%; Pred. No. 1.7e-34;
Matches 94; Conservative 4; Mismatches 5; Indels
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                                                                                                                          63 ESRNEGTIKQAKEKVESKKAEATRLENIKTDRKKAEEEAKRKA 105
                                                                                                                                                              61 ESRNEERIKQAKAKVESKKABATRLEKIKTDRKKABEBAKRKA 103
                                                                                                                                                                                                                                                                                                                               Sequence 19, Application US/09286981B Patent No. 6503511 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 7, Application US/09286981B
; Patent No. 6503511
; GENERAL INFORMATION:
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ORGANISM: Streptococcus pneumoniae
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